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      1489 33.6 1.1 1369462 4 US-09-949-016-13209 Sequence 13209, A
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      C1490 33.8 1.1 390890 4 US-09-949-016-14720 Sequence 14720, A
      C1492 33.6 1.1 158 4 US-09-513-999C-19001 Sequence 19001, A
      C1493 33.6 1.1 278 4 US-09-513-999C-36065 Sequence 36065, A
      C1494 33.6 1.1 345 4 US-08-171-385-20 Sequence 20, App1
      1495 33.6 1.1 345.3 US-08-361-441B-20 Sequence 20, App1
      1496 33.6 1.1 437.3 US-09-265-315E-52 Sequence 52, App1
      1497 33.6 1.1 437.3 US-09-265-315E-52 Sequence 52, App1
      1498 33.6 1.1 437.3 US-09-265-315E-52 Sequence 52, App1
      1499 33.6 1.1 437.4 US-08-361-441B-20 Sequence 52, App1
      1500 33.6 1.1 437.4 US-09-265-315E-52 Sequence 52, App1

ALIGNMENTS
      1
RESULT 1
US-09-489-847-44
Sequence 44, Application US/09489847
Patent No. 6476195
GENERAL INFORMATION:
APPLICANT: Rosen et al
TITLE OR INVENTION: 98 Human Secreted Proteins
FILE REFERENCE: P2031P1
CURRENT FILING DATE: US/09/489, 847
EARLIER FILING DATE: 2000-01-24
EARLIER APPLICATION NUMBER: PCT/US99/17130
EARLIER FILING DATE: 1999-07-29
EARLIER APPLICATION NUMBER: 60/094,657
EARLIER FILING DATE: 1998-07-30
EARLIER APPLICATION NUMBER: 60/095,486
EARLIER FILING DATE: 1998-08-05
EARLIER APPLICATION NUMBER: 60/096,319
EARLIER FILING DATE: 1998-08-12
EARLIER APPLICATION NUMBER: 60/095,454
EARLIER FILING DATE: 1998-08-06
EARLIER APPLICATION NUMBER: 60/095,455
EARLIER FILING DATE: 1998-08-06
NUMBER OF SEQ ID NOS: 376
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 44
LENGTH: 2572
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: SITE
LOCATION: (12527)
OTHER INFORMATION: n equals a,t,g, or c
US-09-489-847-44

Query Match          79.2%; Score 2332.2; DB 4; Length 2572;
Best Local Similarity 99.8%; Pred. No. 0;
Matches 2345; Conservative 0; Mismatches 3; Indels 1; Gaps 1

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QY   CTACTACAAGAATCATTCAGACATCATCCCTGACAGAAAGTGCCCCGAGCTGGGGGAGA 774
Db   CTACTACAAGAATCATTCAGACATCATCCCTGACAGAAAGTGCCCCGAGCTGGGGGAGA 219
QY   TGGAACAATAAAGAAATGCTGAGCTTGTGTGAGCTTTGCTTAATTCTGGCCAACA 834
Db   TGGAACAATAAAGAAATGCTGAGCTTGTGTGAGCTTTGCTTAATTCTGGCCAACA 279
QY   GAAATATCAGCGGCTTAATGTCAACCTCTTGTTCGCCGGGACCTTGAGTGCAGTTGCG 894

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